RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/553,723
Source:	PCT
Date Processed by STIC:	10/28/2005
	7 7

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 10/28/2005 PATENT APPLICATION: US/10/553,723 TIME: 09:17:24

Input Set : A:\PTO.RJ.txt

Output Set: N:\CRF4\10282005\J553723.raw

- 3 <110> APPLICANT: NATIONAL INSTITUTE OF AGROBIOLOGICAL SCIENCES 5 <120> TITLE OF INVENTION: METHOD OF DISTINGUISHING EAR SHAPE AND RESISTANCE AGAINST
- GIBBERELLA ZEAE
 - 6 AND METHOD OF IMPROVING BARLEY PLANT USING THE SAME
 - 8 <130> FILE REFERENCE: MOA-A0301P
- C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/553,723
- C--> 10 <141> CURRENT FILING DATE: 2005-10-14
 - 10 <150> PRIOR APPLICATION NUMBER: JP 2003-110682
 - 11 <151> PRIOR FILING DATE: 2003-04-15
 - 13 <160> NUMBER OF SEQ ID NOS: 7
 - 15 <170> SOFTWARE: PatentIn version 3.1
 - 17 <210> SEQ ID NO: 1
 - 18 <211> LENGTH: 1091
 - 19 <212> TYPE: DNA
 - 20 <213> ORGANISM: Hordeum vulgare
 - 22 <400> SEQUENCE: 1
 - 23 gactgcgtac caattcgctg aggtacttgc tcgcatagtc atggtgctct ttgcaaactg
 25 ccaagaagct ctcgtgcata tagtatgggt ccgcgatgca gatatatttg atgctctccg
 27 tcctgatgaa gtagttgaga tacagcgcat acatgcggat gaactggaaa tccaacttgg
 29 tcatgtggaa catattgtag atgtagtcaa atcgcaggag gaattgattc gcgggccagg
 240
 31 tatcgaagta caccttccca cccggcacct gagccgcgta aagcgggtat cctgaatctt
 300
 - 33 tcacgaccag aaggetttte tetetggeea geacatette atggagtete tttagateat 360
 35 tgtteagtae gtgatetatt gttttggeag gtaggategg ettgeeggeg acatggaaat 420
 - 37 gcttcttacc ttcgacgggt attctgtcta ccacccgggg ttccagaggc ggctagctgc 480
 - 39 ttgaagcate tatggegeet ttaagggeet tteetggeea gtetetteet ttgettettg 540
 41 gtgggtggeg gtagtgaace caccatacet teeetaacea geaceeceag tgteateggg 600
 - 41 gtgggtggcg gtagtgaacc caccatacct teectaacca gcacceccag tgtcatcggg 600
 - 43 ctagacagtg gacggcctcg gggggttgct ggctagaggt ggcatctcga ggcgtctcct 660
 45 gcgaggatgg cttgaagaga gactttttgc attccacctt aggacgttcc ggctccggaa 720
 - 45 gcgaggatgg cttgaagaga gactttttgc attccacctt aggacgttcc ggctccggaa 720 47 aatcaggcaa catcaagtca tcgtctccac actcatagcc tatttcttcg gcgtcatgag 780
 - 49 ccatcaatcc atcataggcg gtattgaaat acttgttcgg gtcctcatca tcatcctcca 840
 - 51 tgtcattatc aacatttgct tcttcgacag tggggtgaca tgatctggca ctaatggagg 900
 - 53 ctctccctcg cgttgtacgc taccatcatc tgccacgaca ggtgcaagta atttggtagg 960
 - 55 tgggatggtg ttcatacctt cttgagaagg tggcgttgga gtgatactgg gctcctcgag 1020 57 acgaataaga gacttcggcc aaggcaagaa tcagttcttg catgcgccaa tcatcttact 1080
 - 59 caggactcat c
 - 62 <210> SEQ ID NO: 2
 - 63 <211> LENGTH: 253
 - 64 <212> TYPE: DNA
 - 65 <213> ORGANISM: Hordeum vulgare
 - 67 <400> SEQUENCE: 2
 - 68 gactgcgtac caattcgact agccatggtt gtgtatgtat ggcacatggg agtaaaacgt 60 70 tacaattctt tttgggaacc acacataatg agtatagcat ggaagatact aataacgttt 120
 - 72 gtcgtaacgt tcacagtaaa gaacaccact caaaattata ttttcaatcc cgctttgaaa 180
 - 74 acttgagete taggaettgt geaaateaat gettacetet geaaagggte tatetatta 240

1091

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76 ctcaggactc atc	253
79 <210> SEQ ID NO: 3	
80 <211> LENGTH: 260	
81 <212> TYPE: DNA	
82 <213> ORGANISM: Hordeum vulgare	
84 <400> SEQUENCE: 3	
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87 gggatgctag ttttgctatg accactacat gattggggta ataatttttc ttattgtgac	120
89 atccccggat traggeraca graatering taattgaact acagtaaaca tatgcaaagg	180
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91 atgccacatc atcgtgattc tattattgat ctcgtgatag tcgaaaccga gtcgaaaatc	260
93 gaagttactc aggactcatc	200
96 <210> SEQ ID NO: 4	
97 <211> LENGTH: 158	
98 <212> TYPE: DNA	
99 <213> ORGANISM: Hordeum vulgare	
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104 ttcctccttc gtgggaggat cgatcttcat caacatattc accagcacca tctcatgtcc	120
106 aaccctagtt catctcgtgt atttactcag gactcatc	158
109 <210> SEQ ID NO: 5	
110 <211> LENGTH: 514	
111 <212> TYPE: DNA	
112 <213> ORGANISM: Hordeum vulgare	
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117 gcacgttcag cttcgtctag acggagtagg caccatcatc atcatgaacc accgtgtcga	120
119 agagateett egagatggtg tegaagaatg teaegeecaa gatgegaeee tateeteaat	180
121 ttggcacgaa ggccttgtca tggatagaag cgcatctcgt cgtgtcgcaa gaatggatat	240
	. 300
123 cgttacaagt acatgtactg aaaagaagag atatatatag aattggctta cactcgccac	360
125 aagctacatc agagtcacat cagtacatta cataatcatc aagagcaaga gcagggtccg	420
127 actacggacg aaaacaaacg ataaaaataa gaacagcgtc cgtccttgct atcccaggct	
129 gccggcctgg aacccatcct agatcgatga agaagaagaa gaagaagcaa ctccaaatga	480
131 acaatcaagg cgctcgcgtt actcaggact catc	514
134 <210> SEQ ID NO: 6	
135 <211> LENGTH: 20	
136 <212> TYPE: DNA	
137 <213> ORGANISM: Artificial	
139 <220> FEATURE:	
140 <223> OTHER INFORMATION: an artificially synthesized primer sequence	e
142 <400> SEQUENCE: 6	
143 atggttgtgt atgtatggca	20
146 <210> SEQ ID NO: 7	
147 <211> LENGTH: 20	
148 <212> TYPE: DNA	
149 <213> ORGANISM: Artificial	
151 <220> FEATURE:	
152 <223> OTHER INFORMATION: an artificially synthesized primer sequence	e
152 <2235 OTHER INFORMATION. An artificially synthesized primer sequence 154 <400> SEQUENCE: 7	_
155 cagaggtaag cattgatttg	20
100 cagaggeaag cacegaceeg	20

RAW SEQUENCE LISTING ERROR SUMMARY

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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,

per 1.823/(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:6,7

VERIFICATION SUMMARY

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date